CRF Errors Corrected by the STIC Systems Branch.    CRF Errors Corrected by the STIC Systems Branch   CRF Processing Dane   C/22/    Changed a file from non-ASCII to ASCII   Changed the margins in cases where the sequence text was wrisepholodown to the next line.    Edited a format error in the Current Application Data section, specifically:     Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other     Added the mandatory heading and subheadings for "Current Application Data".    Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integer	CRF Errors Corrocted by the STIC Systems Branch.    CRF Processing Dame   C/22/    Changed a life from non-ASCII to ASCII   Changed the margins in cases where the sequence text was virsipled down to the next line.    Edited a formal error in the Current Application Data section, specifically:    Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other		May Excel 0590 01/2
Changed a file from non-ASCII to ASCII  Changed the margins in cases where the sequence text was wrisepted down to the next line.  Edited a format error in the Current Application Data section, specifically:  Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other    Added the mandatory heading and subheadings for 'Current Application Data'.  Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integer Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:   non-ASCII 'garbage' at the beginning/end of files;   secretary initials/filename at end of the page numbers throughout text;   other invalid text, such as    Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending step codon in arrino acid sequences and adjusted the '(A)Length,' field accordingly (erroduce to a Patentin bug). Sequences corrected:  Other:	Changed a file from non-ASCII to ASCII  Changed the marginc in cases where the sequence text was wisspital down to the next line.  Edited a format error in the Current Application Data section, specifically:  Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or other    Added the mandatory heading and subheadings for 'Current Application Data'.  Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Defeted extra, invalid, headings used by an applicant, epecifically:  Oeteled:   non-ASCII 'garbage' at the beginning/end of files;   secretary initials/filename at end of the page numbers throughout text.   other invalid text, such as    Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be defeted.  Deleted ending stop coden in argine acid sequences and adjusted the '(A)Length: field accordingly (error due to a Patentin bug). Sequences corrected:  Other:	, N	CRF Errors Corrected by the STIC Systems Branch. CRF Processing Date: 6/22/2
Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other    Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, epecifically:  Deleted:   non-ASCII garbage" at the beginning/end of files;   secretary initials/filename at end of the page numbers throughout text;   other invalid text, such as    Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an origin in the Number of Sequences field, specifically:  A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending stap coden in antino acid sequences and adjusted the "(A)Length: field accordingly (errodue) to a Patentiln bug). Sequences corrected:  Other:	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other    Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, eperfligally:  Life Deleted: non-ASCII garbage at the beginning/end of files: secretary initials/filename at end of the page numbers throughout text; either invalid text, such as subheadings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an orier in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending stop codon in artino acid sequences and adjusted the "(A)Length: field accordingly (errodue to a Patentin bug). Sequences corrected:  Other:	••	
Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other		Changed the margins in cases where the sequence text was wramped down to the next line.
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Deleted:non-ASCII *garbage* at the beginning/end of files: secretary initials/filename at end of	Deleted: non-ASCII *garbage* at the beginning/end of files; secretary initials/filename at end of files; secretary initials		Inserted colons after headings/subheadings. Headings edited included:
page numbers throughout text; other invalid text, such as	page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  eleted ending stop codon in arron acid sequences and adjusted the "(A)Length:" field accordingly (errouge to a Patentin bug). Sequences corrected:		Deleted extra, invalid, headings used by an applicant, specifically: (140), (1417 Alphabetical Leading)
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Deleted <i>endIng</i> stop codon in arrino acid sequences and adjusted the "(A)Length:" field accordingly (errough to a PatentIn bug). Sequences corrected:	Deleted endIng stop codon in artino acid sequences and adjusted the "(A)Length:" field accordingly (errough to a PatentIn bug). Sequences corrected:Other:		
due to a Patentin bug). Sequences corrected:	due to a Patentin bug). Sequences corrected:		A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
Other:	· · · · · · · · · · · · · · · · · · ·	) j:	releted <i>endling</i> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (errolue to a Patentin bug). Sequences corrected:
			Other:

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



OIPE

RAW SEQUENCE LISTING DATE: 06/22/2003 PATENT APPLICATION: US/10/003,632C TIME: 17:43:45

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      8 <130> FILE REFERENCE: CEN0269
     10 <140> CURRENT APPLICATION NUMBER: US/10/003,632C
     11 <141> CURRENT FILING DATE: 2001-11-02
     13 <160> NUMBER OF SEQ ID NOS: 14
     15 <170> SOFTWARE: PatentIn Ver 3.1
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    35 Pro Val Ala Arg Thr Ser Pro Leu Gln Thr Pro Ala Ala Pro Gly Ala
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235

230

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RAW SEQUENCE LISTING DATE: 06/22/2003 PATENT APPLICATION: US/10/003,632C TIME: 17:43:45

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84 Phe Ser Ser Gln Pro Gly His Thr Pro His Pro Ala Ala Ser Arg Asp
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87 Pro Val Ala Arg Thr Ser Pro Leu Gln Thr Pro Ala Ala Pro Gly Ala
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RAW SEQUENCE LISTING DATE: 06/22/2003 PATENT APPLICATION: US/10/003,632C TIME: 17:43:45

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\06202003\J003632C.raw

211 Leu Thr Arg Arg Arg Gly Arg Arg Met Thr Ala Leu Leu Gly Ser Ile 150 155 214 Ala Leu Leu Ala Thr Ile Leu Ala Ala Val Ala Met Ser Arg Arg 215 165 170 217 <210> SEQ ID NO: 6 218 <211> LENGTH: 211 219 <212> TYPE: PRT 220 <213> ORGANISM: Homo sapiens 222 <400> SEQUENCE: 6 223 Met Ala Ser Gly Gln Gly Pro Gly Pro Pro Arg Gln Glu Cys Gly Glu 226 Pro Ala Leu Pro Ser Ala Ser Glu Glu Gln Val Ala Gln Asp Thr Glu 20 25 229 Glu Val Phe Arg Ser Tyr Val Phe Tyr Arg His Gln Gln Glu Gln Glu 40 232 Ala Glu Gly Val Ala Ala Pro Ala Asp Pro Glu Met Val Thr Leu Pro 55 235 Leu Gln Pro Ser Ser Thr Met Gly Gln Val Gly Arg Gln Leu Ala Ile 70 75 238 Ile Gly Asp Asp Ile Asn Arg Arg Tyr Asp Ser Glu Phe Gln Thr Met 85 90 241 Leu Gln His Leu Gln Pro Thr Ala Glu Asn Ala Tyr Glu Tyr Phe Thr 100 105 244 Lys Ile Ala Thr Ser Leu Phe Glu Ser Gly Ile Asn Trp Gly Arg Val 120 247 Val Ala Leu Leu Gly Phe Gly Tyr Arg Leu Ala Leu His Val Tyr Gln 250 His Gly Leu Thr Gly Phe Leu Gly Gln Val Thr Arg Phe Val Val Asp 150 155 253 Phe Met Leu His His Cys Ile Ala Arg Trp Ile Ala Gln Arg Gly Gly 165 170 256 Trp Val Ala Ala Leu Asn Leu Gly Asn Gly Pro Ile Leu Asn Val Leu 185 259 Val Val Leu Gly Val Val Leu Leu Gly Gln Phe Val Val Arg Arg Phe 195 200 262 Phe Lys Ser 263 210 266 <210> SEQ ID NO: 7 267 <211> LENGTH: 170 268 <212> TYPE: PRT 269 <213> ORGANISM: Homo sapiens 271 <400> SEQUENCE: 7 272 Met Ser Gln Ser Asn Arg Glu Leu Val Val Asp Phe Leu Ser Tyr Lys 10 275 Leu Ser Gln Lys Gly Tyr Ser Trp Ser Gln Phe Ser Asp Val Glu Glu 25 278 Asn Arg Thr Glu Ala Pro Glu Gly Thr Glu Ser Glu Met Glu Thr Pro 281 Ser Ala Ile Asn Gly Asn Pro Ser Trp His Leu Ala Asp Ser Pro Ala

RAW SEQUENCE LISTING DATE: 06/22/2003 PATENT APPLICATION: US/10/003,632C TIME: 17:43:45

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\06202003\J003632C.raw

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284 Val Asn Gly Ala Thr Gly His Ser Ser Ser Leu Asp Ala Arg Glu Val
                        70
287 Ile Pro Met Ala Ala Val Lys Gln Ala Leu Arg Glu Ala Gly Asp Glu
                    8.5
                                        90
290 Phe Glu Leu Arg Tyr Arg Arg Ala Phe Ser Asp Leu Thr Ser Gln Leu
               100
293 His Ile Thr Pro Gly Thr Ala Tyr Gln Ser Phe Glu Gln Asp Thr Phe
                                120
296 Val Glu Leu Tyr Gly Asn Asn Ala Ala Ala Glu Ser Arg Lys Gly Gln
299 Glu Arg Phe Asn Arg Trp Phe Leu Thr Gly Met Thr Val Ala Gly Val
300 145
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302 Val Leu Leu Gly Ser Leu Phe Ser Arg Lys
303
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305 <210> SEQ ID NO: 8
306 <211> LENGTH: 160
307 <212> TYPE: PRT
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314 Leu Tyr Glu Gln Leu Leu Glu Pro Pro Thr Met Glu Val Leu Gly Met
                                    25
317 Thr Asp Ser Glu Glu Asp Leu Asp Pro Met Glu Asp Phe Asp Ser Leu
320 Glu Cys Met Glu Gly Ser Asp Ala Leu Ala Leu Arg Leu Ala Cys Ile
                            55
323 Gly Asp Glu Met Asp Val Ser Leu Arg Ala Pro Arg Leu Ala Gln Leu
                        70
326 Ser Glu Val Ala Met His Ser Leu Gly Leu Ala Phe Ile Tyr Asp Gln
                                        90
329 Thr Glu Asp Ile Arg Asp Val Leu Arg Ser Phe Met Asp Gly Phe Thr
               100
                                    105
332 Thr Leu Lys Glu Asn Ile Met Arg Phe Trp Arg Ser Pro Asn Pro Gly
    115
                                120
                                                    125
335 Ser Trp Val Ser Cys Glu Gln Val Leu Leu Ala Leu Leu Leu Leu
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338 Ala Leu Leu Pro Leu Leu Ser Gly Gly Leu His Leu Leu Lys
339 145
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341 <210> SEQ ID NO: 9
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343 <212> TYPE: PRT
344 <213> ORGANISM: Homo sapiens
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350 Glu Gln Ile Met Lys Thr Gly Ala Leu Leu Leu Gln Gly Phe Ile Gln
               20
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/22/2003 PATENT APPLICATION: US/10/003,632C TIME: 17:43:46

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\06202003\J003632C.raw

## Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

VERIFICATION SUMMARY

DATE: 06/22/2003 PATENT APPLICATION: US/10/003,632C TIME: 17:43:46

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\06202003\J003632C.raw

L:614 M:283 W: Missing Blank Line separator, <400> field identifier

10/003,6320

## Sequence Listing

<110> Lee, Chichang; Ly, Celia; Moore, Gordon; Chi, Xiamei

<120> Methods and Compositions for Enhanced Protein Expression and/or Growth of Cultured Cells Using Co-Transcription of a Bcl2 Encoding Nucleic Acid

<130> CEN0269

<140> CHURENT APPLICATION NUMBER! US/10/003,6328 <141> CURRENT APPLICATION DATE! 2001-11-02

> Does Nor Comply Corrected Diskette Needed